

# ChIP coupled to quantitative RT-PCR (qRT-PCR)

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 An abbreviated version of this protocol was published in eLIFE in Oct 2018

Chromatin mapping identifies BasR, a key regulator of bacteria-triggered production of fungal secondary metabolites

DOI: 10.7554/eLife.40969

## Detailed protocol

Dear colleague, the protocol is published, see attached file,  
all the best  
Joseph

## Related files

 Boedi\_MethMolBiol\_ChIP.pdf



**How to cite:** (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Strauss, J. (2020). ChIP coupled to quantitative RT-PCR (qRT-PCR). Bio-protocol Preprint. [bio-protocol.org/prep378](https://bio-protocol.org/prep378).
2. Fischer, J., Müller, S. Y., Netzker, T., Jäger, N., Gacek-Matthews, A., Scherlach, K., Stroe, M. C., García-Altares, M., Pezzini, F., Schoeler, H., Reichelt, M., Gershenzon, J., Krespach, M. K., Shelest, E., Schroeckh, V., Valiante, V., Heinzel, T., Hertweck, C., Strauss, J. and Brakhage, A. A. (2018). Chromatin mapping identifies BasR, a key regulator of bacteria-triggered production of fungal secondary metabolites. eLIFE. DOI: [10.7554/eLife.40969](https://doi.org/10.7554/eLife.40969)

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